

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/572,882  
Source: IFWP  
Date Processed by STIC: 3/31/06

***ENTERED***



IFWP

## RAW SEQUENCE LISTING

DATE: 03/31/2006

PATENT APPLICATION: US/10/572,882

TIME: 12:19:12

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\03292006\J572882.raw

3 <110> APPLICANT: Bayer HealthCare AG  
 4 Golz, Stefan  
 5 Broggemeier, Ulf  
 6 Geerts, Andreas  
 8 <120> TITLE OF INVENTION: Diagnostics and Therapeutics for Diseases  
 Associated with  
 9 G-Protein Coupled Receptor ADIPOR2 (ADIPOR2)  
 11 <130> FILE REFERENCE: LeA.36 902  
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/572,882  
 C--> 13 <141> CURRENT FILING DATE: 2006-03-20  
 13 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/010383  
 14 <151> PRIOR FILING DATE: 2004-09-16  
 16 <150> PRIOR APPLICATION NUMBER: EP03021898.6  
 17 <151> PRIOR FILING DATE: 2003-09-27  
 19 <160> NUMBER OF SEQ ID NOS: 5  
 21 <170> SOFTWARE: PatentIn version 3.3  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 3500  
 25 <212> TYPE: DNA  
 26 <213> ORGANISM: Homo sapiens  
 28 <400> SEQUENCE: 1  
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 31 agactggctc aaggataatg acttctctt gcatggacac cggcctccta tgccttcttt 120  
 33 ccgggctgtg tttaagagca ttttcagaat acacacagaa acaggcaaca ttggacaca 180  
 35 tctcttaggt tgtgtatcct tcctgtgcct ggggatcttt tatatgtttc gcccaaatat 240  
 37 ctccctttgtg gcccctctgc aagagaaggt ggtctttgga ttatttttct taggagccat 300  
 39 tctctgcctt tctttttcat ggctcttcca cacagtctac tgccactcag agggggtctc 360  
 41 tcggctcttc tctaaactgg attactctgg tattgctctt ctgattatgg gaagttttgt 420  
 43 tccttggctt tattattctt tctactgtaa tccacaacct tgcttcatct acttgattgt 480  
 45 catctgtgtg ctgggcattg cagccattat agtctcccag tgggacatgt ttgccacccc 540  
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 49 taccttgcat tatgtcatct cggaggggtt ccttaaggcc gccaccatag ggcagatagg 660  
 51 ctggttgatg ctgatggcca gcctctacat cacaggagct gccctgtatg ctgcccggt 720  
 53 ccccgaaacgc tttttccctg gcaaatgtga catctgggtt cactctcatc agctgtttca 780  
 55 tatctttgtg gttgctggag cttttgttca cttccatggt gtctcaaacc tccaggagtt 840  
 57 tcgtttcatg atcggcgagg gctgcagtga agaggatgca ctgtgatacc taccagtctc 900  
 59 cagggactat gaccctaaac cagggcctgc ggcacttgcg ggctccctg ctggctactg 960  
 61 atgccagtac cagaggagcc ccaaaacttt gacagcctcg tgggctttgt gacggcccag 1020  
 63 gggctctgcg tggtagatga ctgagaagag aaaaacaaaa ataaatcata cctcaaagga 1080  
 65 tggagtgcac caattgggag aaaaggagac atagcccaaa ccctggctta ttcttgggat 1140  
 67 ctactgattg cgggctctgc aagacccttg gcaaaactggc ttctgatcca tatcatattt 1200  
 69 attttagaaa gatggcgaaa cagttagct ggtggttctt tcttctccct ttctctctct 1260  
 71 ctatgacaat aatacaaac aatttaagt aacatttata tccgataagg ggtgggagt 1320  
 73 tgattttaaa tgctcttttg ggagaacaaa gaaattaatg taaataagat ttctaactgt 1380

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79 taggatcagg tgatagcccc ggaatgtaca gtgtcttggg gcaccaagat gccttctaaa 1560
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89 ctagaacat tagcatgggc tgatctgatt acttcctggc atcccgtca cctttatggg 1860
91 aagtcttatt agagggatgg gacagttttc catatccttg ctgtggagct ctggaacact 1920
93 ctctaaattt ccctctatta aaaatcactg ccctaactat acttcctcct tgagggaata 1980
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127 acagaagtag cagcctctgg gtgcagtcac ccacacccca aagctggaag gatctggttc 3000
129 aacatagcac aaacccttag gaaaaatgaa attaacatca ctgatgtgta atccagtaaa 3060
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137 tttttaattt tcttttttgg ccctaggctg gttgggacct ctacagcttc attctttcac 3300
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141 acaaagcaca tttttgggga tcatagaagg ttggggttcc agaaaggcat ctgtgtgatg 3420
143 gttccattga tgtgggattt ccctacttgc tgtattctca gtttctaata aaaagaacca 3480
145 aatgaaaaaa aaaaaaaaaa 3500
148 <210> SEQ ID NO: 2
149 <211> LENGTH: 258
150 <212> TYPE: PRT
151 <213> ORGANISM: Homo sapiens
153 <400> SEQUENCE: 2
155 Met Pro Ser Phe Arg Ala Cys Phe Lys Ser Ile Phe Arg Ile His Thr
156 1 5 10 15
159 Glu Thr Gly Asn Ile Trp Thr His Leu Leu Gly Cys Val Phe Phe Leu
160 20 25 30
163 Cys Leu Gly Ile Phe Tyr Met Phe Arg Pro Asn Ile Ser Phe Val Ala
164 35 40 45
167 Pro Leu Gln Glu Lys Val Val Phe Gly Leu Phe Phe Leu Gly Ala Ile
168 50 55 60

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171 Leu Cys Leu Ser Phe Ser Trp Leu Phe His Thr Val Tyr Cys His Ser
172 65                      70                      75                      80
175 Glu Gly Val Ser Arg Leu Phe Ser Lys Leu Asp Tyr Ser Gly Ile Ala
176                      85                      90                      95
179 Leu Leu Ile Met Gly Ser Phe Val Pro Trp Leu Tyr Tyr Ser Phe Tyr
180                      100                     105                     110
183 Cys Asn Pro Gln Pro Cys Phe Ile Tyr Leu Ile Val Ile Cys Val Leu
184                      115                     120                     125
187 Gly Ile Ala Ala Ile Ile Val Ser Gln Trp Asp Met Phe Ala Thr Pro
188                      130                     135                     140
191 Gln Tyr Arg Gly Val Arg Ala Gly Val Phe Leu Gly Leu Gly Leu Ser
192 145                      150                      155                      160
195 Gly Ile Ile Pro Thr Leu His Tyr Val Ile Ser Glu Gly Phe Leu Lys
196                      165                      170                      175
199 Ala Ala Thr Ile Gly Gln Ile Gly Trp Leu Met Leu Met Ala Ser Leu
200                      180                      185                      190
203 Tyr Ile Thr Gly Ala Ala Leu Tyr Ala Ala Arg Ile Pro Glu Arg Phe
204                      195                      200                      205
207 Phe Pro Gly Lys Cys Asp Ile Trp Phe His Ser His Gln Leu Phe His
208                      210                      215                      220
211 Ile Phe Val Val Ala Gly Ala Phe Val His Phe His Gly Val Ser Asn
212 225                      230                      235                      240
215 Leu Gln Glu Phe Arg Phe Met Ile Gly Gly Gly Cys Ser Glu Glu Asp
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219 Ala Leu

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223 &lt;210&gt; SEQ ID NO: 3

224 &lt;211&gt; LENGTH: 20

225 &lt;212&gt; TYPE: DNA

226 &lt;213&gt; ORGANISM: Artificial

228 &lt;220&gt; FEATURE:

229 &lt;223&gt; OTHER INFORMATION: Primer1 (forward primer)

231 &lt;400&gt; SEQUENCE: 3

232 catggtgtct caaacctcca

20

235 &lt;210&gt; SEQ ID NO: 4

236 &lt;211&gt; LENGTH: 20

237 &lt;212&gt; TYPE: DNA

238 &lt;213&gt; ORGANISM: Artificial

240 &lt;220&gt; FEATURE:

241 &lt;223&gt; OTHER INFORMATION: Primer2 (reverse primer)

243 &lt;400&gt; SEQUENCE: 4

244 cagtgcattcc tcttcactgc

20

247 &lt;210&gt; SEQ ID NO: 5

248 &lt;211&gt; LENGTH: 23

249 &lt;212&gt; TYPE: DNA

250 &lt;213&gt; ORGANISM: Artificial

252 &lt;220&gt; FEATURE:

253 &lt;223&gt; OTHER INFORMATION: Probe1

255 &lt;400&gt; SEQUENCE: 5

256 agtttcgttt catgatcggc ggg

23

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/572,882

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Input Set : A:\PTO.KD.txt  
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/572,882

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Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\03292006\J572882.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date